

AUG 0 1 2002

TECH CENTER 1600/2900



1600

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/952,570B

DATE: 07/23/2002

TIME: 14:37:20

Input Set : A:\08269-006001.txt

Output Set: N:\CRF3\07232002\H952570B.raw

ENTERED

4 <110> APPLICANT: Lunderbert, Joakim Uhlen, Mathias 7 <120> TITLE OF INVENTION: METHOD FOR ACTIVATING A REVERSIBLY INACTIVATED IMMOBILIZED ENZYME BY RELAEASE FROM AN IMMOBILIZING MOIETY AND ITS USE IN NUCLEIC ACID 9 AMPLIFICATION REACTIONS 11 <130> FILE REFERENCE: 08269-006001 13 <140> CURRENT APPLICATION NUMBER: US 08/952,570B C--> 14 <141> CURRENT FILING DATE: 1997-11-10 16 <150> PRIOR APPLICATION NUMBER: PCT/GB96/01108 17 <151> PRIOR FILING DATE: 1996-05-09 19 <150> PRIOR APPLICATION NUMBER: GB/9509336.5 20 <151> PRIOR FILING DATE: 1995-05-09 22 <160> NUMBER OF SEQ ID NOS: 23 24 <170> SOFTWARE: FastSEQ for Windows Version 4.0 26 <210> SEQ ID NO: 1 27 <211> LENGTH: 32 28 <212> TYPE: DNA 29 <213> ORGANISM: Thermus aquaticus 31 <400> SEQUENCE: 1 32 gggaattcca tggacgatct gaagctctcc tg 32 34 <210> SEQ ID NO: 2 35 <211> LENGTH: 32 36 <212> TYPE: DNA 37 <213> ORGANISM: Thermus aquaticus 39 <400> SEQUENCE: 2 32 40 ccccaagett etgeaggate ttetecaega tg 42 <210> SEQ ID NO: 3 43 <211> LENGTH: 20 44 <212> TYPE: DNA 45 <213> ORGANISM: Thermus aquaticus 47 <400> SEQUENCE: 3 20 48 gaagatcctg cagtaccggg 50 <210> SEQ ID NO: 4 51 <211> LENGTH: 26 52 <212> TYPE: DNA 53 <213> ORGANISM: Thermus aquaticus 55 <400> SEQUENCE: 4 56 gggtcgacct ccttggcgga gagcca 26 58 <210> SEQ ID NO: 5 59 <211> LENGTH: 15 60 <212> TYPE: DNA 61 <213> ORGANISM: Thermus aquaticus 63 <400> SEQUENCE: 5

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Input Set : A:\08269-006001.txt

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RAW SEQUENCE LISTING

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Input Set : A:\08269-006001.txt

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/23/2002 PATENT APPLICATION: US/08/952,570B TIME: 14:37:21

Input Set : A:\08269-006001.txt

Output Set: N:\CRF3\07232002\H952570B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 1 Seq#:13; N Pos. 1 Seq#:14; N Pos. 1 Seq#:15; N Pos. 1 Seq#:16; N Pos. 1 Seq#:20; N Pos. 1